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Data Visualization

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The Data as a Quantity

- Quantities can be classified in two categories:
 - Intrinsically continuous (scientific visualization, or scivis)
 - e.g. pressure, temperature, position, speed, density, force, color, light intensity etc.
 - Intrinsically discrete (information visualization, or infovis)
 - e.g. text, hypertext, content of web pages, database records etc.
- Sampled data originally continuous data represented in a finite approximative form
- Corollary of the difference between sampled and discrete data:

In the case of sampled data, we can go back to a continuous approximation of the original (intrinsically) continuous data but it make no sense for (intrinsically) discrete data

Continuous Data

• Continuous data as a function:

$$f: D \to C$$

• Intuitive interpretation of continuity:



Source: Data Visualization: Principles and Practice

Datasets and Dimensions

- Let the triplet D = (D, C, f) define a continuous dataset
- We assume that $f: D \to C$
- *D* refers to a function domain, *C* is the function codomain
- d is the geometrical dimension of the space \mathbb{R}^d into which D is embedded
- s represents the topological dimension of D itself (e.g. plane in the Euclidean space \mathbb{R}^3 has s = 2)
 - It holds that $s \leq d$
 - *s* is number of independent variables required to represent the domain *D*
- Codimension of an object of some d and s is the difference d-s

Datasets and Dimensions

- Virtually all data-visualization applications fix geometrical dimension to d = 3
- Only the topological dimension varies $s = \{1, 2, 3\}$
 - s = 1 corresponds to curves
 - s = 2 corresponds to surfaces
 - s = 3 corresponds to the volumetric datasets
- Topological dimension and dataset dimension are often used interchangeably
- Topological dimension is important in case of sampled datasets and when choosing a grid cell type

Datasets and Dimensions

- Function values are called dataset attributes
- The dimensionality *c* of the function codomain *C* is also called the attribute dimension
- Typically ranges from 1 to 4
- E.g. temperature assigned to some point in the Euclidean space \mathbb{R}^3 has c = 1 (it is a scalar value)

- Typically, continuous functional representation of data is not available
- Moreover, several operations (filtering, simplification, analysis etc.) on continous data are not efficient
- Visualization applications work predominantly with sampled datasets
- Important operations relating continuous and sampled data:
 - Sampling quite straightforward
 - Reconstruction more complicated, the goal is to recover an approximated version of the original continuous data

- Reconstruction employs interpolation of the values of the function between its sample points
- Two basic forms of sampling strategies:
 - Uniform
 - Non-uniform (e.g. respecting the distribution of the importance of the sampled data)
- Sampled dataset should be accurate (up to an user-specified error), minimal (w.r.t. an error), generic (operations), efficient (algorithmically), and simple (implementation) [Schroeder et al. 2006]

- Sampled dataset $\{f_i, p_i\}$ consists of a set of N sample points and values
- Interpolation the reconstructed function should equal the original one at all sample points, i.e. $\tilde{f}(p_i) = f(p_i) = f_i$ Basis/interpolation functions
- One way to define reconstruction function: $\tilde{f} = \sum_{i=1}^{N} f_i \dot{\phi}_i$
- Subsequently, we get $\tilde{f}(p_j) = \sum_{i=1}^N f_i \phi_i(p_j) = f(p_j) = f_j$ for $\forall j$
- Orthogonality basis function $\phi_i(p_j) = \begin{cases} 1, & i = j \\ 0, & i \neq j \end{cases}$ • Normality of basis function $\sum_{i=1}^N \phi_i(x) = 1, \forall x \in D$



Domain Subdivision

- A grid (aka mesh) is a subdivision of a domain *D* into a collection of cells (aka elements)
- Most commonly used cells:
 - Polylines in $\mathbb R$
 - Polygons in \mathbb{R}^2
 - Polyhedra in \mathbb{R}^3
- Union of cells cover entire domain *D* and cells are non-overlapping, and vertices are sample points



Domain Subdivision

- The most common domain discretization used in SciVis are simplicial complexes
- A d-simplex is the convex hull of d + 1 affinely independent points in \mathbb{R}^d is d-dimensional
- A simplicial complex is a finite collection of simplices that contains all faces of any simplex and where the intersection of two simplices empty or a face of both



Constant Basis Function

constant, zero-order continuity global basis function

• Simplest set of basis function:

$$x) = \begin{cases} 1, & x \in c_i \\ 0, & x \notin c_i \end{cases}$$

- Sample points are inside the grid cells
- Nearest-neighbor interpolation
- Virtually no computation cost
- Work with any cell shape and in any dimension
- Provide a poor, staircase-like approximation
- We can provide a better (i.e. more continuous) reconstruction of the original function

Linear Basis Functions

- Linear basis function next simplest basis functions
- Need to make some assumption about the cell types used in the grid
- Assume quadrilateral cells having 4 vertices
- Reference quad cell in \mathbb{R}^2 : $v_1 = (0,0)$, $v_2 = (1,0)$, $v_3 = (1,1)$, $v_4 = (0,1)$

Set of local basis functions

Barycentric Coordinates

- A simplex is a convex hull of k + 1 points in a k-dimensional space
- Barycentric coordinates provide a simple way to interpolate over simplices
- In case of (planar) triangles, k = 2



From this equation is clear what the barycentric coordinates r and s actually mean

	$p1 \coloneqq \begin{bmatrix} 27.2\\11.8 \end{bmatrix} \qquad p$	$D2 \coloneqq \begin{bmatrix} 68.4\\ 16.3 \end{bmatrix}$	$p3 \coloneqq \begin{bmatrix} 46.6\\40.9 \end{bmatrix}$		
	A1:=232.2	42 := 186.24	$A3 \coloneqq 137.37$		
	$A \coloneqq A1 + A2 + A3 = 555.81$				
	$r\!\coloneqq\!\frac{A1}{A}\!=\!0.418$	$s \coloneqq \frac{A2}{A} = 0.33$	$5 \qquad t \coloneqq \frac{A3}{A} = 0.247$		
	$r\!+\!s\!+\!t\!=\!1$		1 - r - s = 0.247		
	$p \coloneqq r \cdot p1 + s \cdot p2 + t \cdot p3 = \begin{bmatrix} 45.8\\ 20.5 \end{bmatrix}$				
	$p \coloneqq r \cdot p1 + s \cdot p2 + (1 - r - s) \cdot p3 = \begin{bmatrix} 45.8\\20.5 \end{bmatrix}$ or in the rearranged form				
	$p \coloneqq r \cdot p1 + s \cdot p2 + 1 \cdot p3 - r \cdot p3 - s \cdot p3 = \begin{bmatrix} 45.8\\ 20.5 \end{bmatrix}$				
	$p \coloneqq r \cdot (p1 - p3) + s$	$s \cdot (p2 - p3) + 1 \cdot$	$p_3 = \begin{vmatrix} 45.8\\20.5 \end{vmatrix}$		

Barycentric Coordinates for Triangles

• Barycentric (area) coordinates [r, s, t] describe location of a point p in a triangle in relation to vertices v_i

$$p = [r, s, t] = r v_1 + s v_2 + t v_3$$
,

where $r, s, t \ge 0$ and r + s + t = 1

- Coordinates corresponds to the signed area of the opposite subtriangle divided by area of the triangle
- Note that the point *p* is uniquely defined by any two of the three barycentric coordinates, e.g. *r* and *s*

$$p = [r,s] = r v_1 + s v_2 + (1 - r - s) v_3 \text{ or } T(r,s) = \sum_{i=1}^3 v_i \Phi_i^1(r,s)$$

• In the same way, we can interpolate any quantity (or function) inside the triangle

Forward Transformations

• Given any cell type having n vertices p_i in \mathbb{R}^3 , we define transformation T that maps from a point [r, s] in reference cell coordinate system to a point [x, y, z] in the actual cell as follows

$$p = [x, y, z] = T(r, s) = \sum_{i=1}^{n} p_i \Phi_i^1(r, s)$$

- *T* maps the reference cell to the world cell
- T^{-1} maps points [x, y, z] in the world cell to points [r, s] in the reference cell

Backward Transformation

• Having T^{-1} , we can rewrite the reconstruction function $\tilde{f} = \sum_{i=1}^{n} f_i \Phi_i$ for quad cell as $\tilde{f}(x, y, z) = \sum_{i=1}^{4} f_i \Phi_i^1 (T^{-1}(x, y, z))$

• To compute T^{-1} , we have to invert the expression $T(r,s) = \sum p_i \Phi_i^1(r,s)$

• Given a rectangular cell, this yields

$$T_{\rm rect}^{-1}(x,y,z) = (r,s) = \left(\frac{(p-p_1)\cdot(p_2-p_1)}{\|p_2-p_1\|^2}, \frac{(p-p_1)\cdot(p_4-p_1)}{\|p_4-p_1\|^2}\right)$$

 Now, we have a simple way how to reconstruct a piecewise C¹ function from samples on any rectangular grid. Arbitrary quad cells require some more elaborated numerical solution for obtaining r, s

Backward Transformation for Triangles

- Same situation but with triangle cell (simplest cell in 2D)
- Three linear basis functions

$$\Phi_{1}^{1}(r,s) = r$$

$$\Phi_{2}^{1}(r,s) = s$$

$$\Phi_{3}^{1}(r,s) = 1 - r - s$$

• The transformation T^{-1} for triangular cells

$$T_{\rm tri}^{-1}(x,y,z) = (r,s) = \left(\frac{\|(p-p_1) \times (p_3-p_1)\|}{\|(p_2-p_1) \times (p_3-p_1)\|}, \frac{\|(p-p_1) \times (p_2-p_1)\|}{\|(p_3-p_1) \times (p_2-p_1)\|}\right)$$

Cells







Source: Data Visualization: Principles and Practice



Source: Data Visualization: Principles and Practice

Rectangular Cell

$$\begin{split} \Phi_1^1(r,s) &= (1-r)(1-s), \\ \Phi_2^1(r,s) &= r(1-s), \\ \Phi_3^1(r,s) &= rs, \\ \Phi_4^1(r,s) &= (1-r)s. \end{split}$$



$$T_{\rm rect}^{-1}(x,y,z) = (r,s) = \left(\frac{(p-p_1)\cdot(p_2-p_1)}{\|p_2-p_1\|^2}, \frac{(p-p_1)\cdot(p_4-p_1)}{\|p_4-p_1\|^2}\right)$$

$$\begin{split} \Phi^1_1(r,s) &= (1-r)(1-s), \\ \Phi^1_2(r,s) &= r(1-s), \\ \Phi^1_3(r,s) &= rs, \\ \Phi^1_4(r,s) &= (1-r)s. \end{split}$$

• The world coordinates of a point p on the given quadrilateral cell with known parametric coordinates r, s can be computed as follows

$$p = (x, y, z)^T = T_{quad}(r, s) = \sum_{i=0}^4 p_i \Phi_i^1(r, s)$$
,

where T_{quad} is a simple bilinear interpolation on a rectangle

$$T_{quad}(r, s) = (1 - s)[(1 - r)p_1 + rp_2] + s[rp_3 + (1 - r)p_4]$$

• Typically, we would like to obtain interpolated quantity f at this point

$$\tilde{f}(x, y, z) = \sum_{i=1}^{4} f_i \Phi_i^1(T_{quad}^{-1}(x, y, z))$$

Unlike in previous cases, we cannot guess inverse of T_{quad} directly.



We are looking for the interpolated valued of f at the point p and to do so, we need to find the corresponding point q in reference coordinates to be able to evaluate $\tilde{f}(p)$.

To find the zeroes of a single vector-valued function $F: \mathbb{R}^k \to \mathbb{R}^{k \text{ or } m}$ we may use Newton's method: $x_{n+1} = x_n - J_F(x_n)^{-1}F(x_n)$

• One solution is to numerically solve for r, s as functions of x, y(z) by Newton's method (note that $T_{quad} = T: \mathbb{R}^2 \to \mathbb{R}^3$ here)

$$\binom{r^{t+1}}{s^{t+1}} = \binom{r^t}{s^t} - \boldsymbol{J}_T^{-1}(r^t, s^t)(\boldsymbol{T}(r, s) - \boldsymbol{p})$$

where J_T^{-1} is the generalized inverse of the non-square Jacobian matrix

Also note that matrix J_T evolves over time as well as r and s get updated during iterations

$$\boldsymbol{J}_{T}(r,s) = \begin{pmatrix} \frac{\partial T}{\partial r} & \vdots & \frac{\partial T}{\partial s} & \frac{\partial T}{\partial s} \\ \vdots & \vdots & \vdots \end{pmatrix}_{2}$$

Matrix describes direction and speed of position changes of T when r, s are varied.

2 or 3×2

• It is easy to see that

$$\frac{\partial \boldsymbol{T}}{\partial r}(r,s) = (s-1)(\boldsymbol{p}_1 - \boldsymbol{p}_2) + s(\boldsymbol{p}_3 - \boldsymbol{p}_4)$$

and

$$\frac{\partial \boldsymbol{T}}{\partial s}(r,s) = (r-1)(\boldsymbol{p}_1 - \boldsymbol{p}_4) + r(\boldsymbol{p}_3 - \boldsymbol{p}_2)$$

The pseudo inverse of J_T can be computed by function
 cv::invert(J, J_inv, cv::DECOMP_SVD); // C++ (but it is terribly slow)
 J_inv = numpy.linalg.pinv(J) # Python

Example on General Quad Cell

• Test example for the afore described procedure:

For the cell with vertices
$$p_1 = \begin{pmatrix} 1 \\ 0 \\ 1 \end{pmatrix}$$
, $p_2 = \begin{pmatrix} 3 \\ 0.25 \\ 1 \end{pmatrix}$, $p_3 = \begin{pmatrix} 4 \\ 3 \\ 1 \end{pmatrix}$, $p_4 = \begin{pmatrix} 0 \\ 3.5 \\ 1 \end{pmatrix}$ and the query point $p = \begin{pmatrix} 1.8 \\ 2.7 \\ 1 \end{pmatrix}$ we get $r = 0.445$ and $s = 0.818$ just after 3 iterations while the initial estimates of r^0 and s^0 are set to 0.5.

 If you get the same r and s for given vertices, your implementation is probably correct

Exercise

- Typically, finite-element meshes are generated from constructive-solid-geometry (CSG) models during finite element analysis (FEA) used in engineering
- Triangular and quadrilateral subdivisions of simulation domains are the most

